

FIGURE 1

Human Basic Fibroblast Growth Factor

1
 AAT TCA TGC CTC TTT CTC TCC TTT TGT TGG TAG AGC ACT TCA GCC TCT GTC CTT 37
 TAA TTT TAA AGT TTA TGC CCC ACT TGT ACC CCT GGT TTT TGG GTG ATT TAG AAG 61
 TTT TCA AAG CCT GCT CTG ACA CAG ACT CTT GCT TGG ATT GCA ACT TCT CTA CTT 125
 TGG GGT GGA AAC GGC TTC TCC GTT TTT AAA CCG TAG CCG GGA AAA AAT GGG GGA 149
 CAA AGT TGA GTT TAA ACT TTT AAA AGT TGA GTC AGC GCT GGT TGC GCA GGA AAA 243
 GCC CCG CAG TGT GGA GAA AGC CTA AAC GTG GTT TGG GTG GTG CCG GGG TTG GGC 267
 GGG GGT GAC TTT TGG GGG ATA AAG GGC GGT GGA GCC CAG GGA ATG CCA AAG CCC 331
 TGC GGC GGC CTC GCA CCG GCG CCC CCC GCC CTT CCG CTC TCC CCG GCG CCG GAC 455
 TGA GGC CCG GCT CCC CCG CCG ACT GAT GTC GCG CCG TTT GGT GTT GTC GCG GAA 459
 GCG GCC GAA CTC AGA GCG CCG CCC CAG AAA ACT CCA GCG AGT AGC GCG CCG CCG 513
 GCA GGA GGG AGG AGA ACT GGG GCG GCG GGA GCG TGG TGG GTG TGG GGG GTG GAG 567
 ATG TAG AAG ATG TGA CCG CCG GGC CCG GCG GGT GCG AGA TTA GCG GAC GCG TGC 621
 CCG CCG TTT CAA CCG GAT CCC GGG CCG TGC AGC TTT GGA GCG GCG TCT CCC CAG 675
 GCG GCG TCG GCG GAG AGA CCC ATC TGT GAA CCC CAG GTC CCG GCG CCG CCG CTC 729
 GCC GCG CAC CAG GCG CCG GCG GAG AGA GCG GCG GCG GCG GCG GCG GCG GCG GCG 783
 GGA CCG CCG GCG CCG CCG CCG GCT GCT GCG CCG GCG GCG GCG GCG GCG GCG GCG 837
 GGG CCG TCG CCG GAG CCG GTT GGA GCG CCG GCG CCG GCG GCG GCG GCG GCG GCT 891
 CCG GCG GCG GCT CCA GCG GCT CCG GGA TCC CCG CCG CCG CCG GCA GCG ACC ATG 945
 ALE Gly Ser Ile Thr Thr Thr Phe Ala Leu Phe Phe Gln Asp Gly Gly Ser Gly 1026
 GCG TTC CCG GCG GCG CAC TTC AAG CAG CCC AAG CCG CTG TAC TGC AAA AAG GCG 1051
 ALE Phe Phe Phe Gly Ile Phe Lys Asp Phe Lys Arg Leu Tyr Cys Lys Asn Gly 1088
 GCG TTC TTC CTG GCG ATC CAG CCG CCG GCG GCG GCG GCG GCG GCG GCG GCG GCG 1126
 Gly Phe Phe Leu Arg Ile Ile Phe Asp Gly Arg Val Asp Gly Val Arg Gln Lys 1161
 ACG GAC CCG CAC ATC AAG CTA CCA CTA CCA GCA GAA CAG ACA GGA GTT GTG TCT 1190
 Ser Asp Phe Phe Ile Ile Lys Leu Gln Leu Gln Ala Gln Gln Arg Gly Val Val Ser 1215
 ATC AAG GAA GTG TGT GCT AAC CTT TAC CTT GCT ATG AAG GAA GAT GGA AGA TTA 1269
 Ile Lys Gly Val Cys Ala Asn Arg Tyr Leu Ala Asn Lys Gln Asp Gly Arg Leu 1311
 CTT GCT CTT AAA TGT GTT ACG GAT GAG TGT TTT TTT GAA CCA TTA GAA TCT 1369
 Leu Ala Ser Lys Cys Val Thr Asp Gln Cys Phe Phe Gln Arg Leu Gln Ser 1408
 AAT AAG TAC AAT ACT TAC CCG TCA AGC AAA TAC ACC AGT TGG TAT GTG GCA TTT 1431
 Asn Asn Tyr Asn Thr Tyr Arg Ser Arg Lys Tyr Thr Ser Trp Tyr Val Ala Leu 1464
 AAA CCA ACT GCG CAG TAT AAA CTT GCA TCC AAA ACA GGA CTT GCG CAG AAA GCT 1477
 Lys Arg Thr Gly Gln Lys Tyr Lys Leu Gly Ser Lys Thr Gly Phe Gly Gln Lys Ala 1519
 ATA TTT TTT CTT CCA ATG TGT GCT AAG AGC TGA TTT TAA TGG CCA CAT CTA ATC 1555
 ALE Phe Leu Phe Phe Phe Ser Ser Ala Lys Ser 1585
 TCA TTT CAC ATG AAA GAA GAA GTA TAT TTT AGA AAT TTT TTA ATG AGA GTA AAA 1621
 SAA AAT AAA TGT GTA TAG CTC AGT TTT GAT AAT TGG TCA AAC AAT TTT TTA TCC 1675
 AGT AGT AAA ATA TGT AAC CAT GCG CAG TAA AGA AAA ATA ACA AAA GTT GTA AAA 1701
 TGT ATA TTC TCC CTT TTA TAT TGC ATC TGG TGT TAC CCA GTG AAG CTT ACC TAG 1755
 ACG AAT GAT CTT TTT CAC GCA TCT GCT TTA TTC GAA AAG AGG CTT TTA AAA TGT 1783
 GCA TGT TTA GAA AAC AAA ATT TCT TCA TGG AAA TCA TAT ACA TTA GAA AAT CAC 1837
 AGT CAG ATG TTT AAT CAA TCG AAA AAT GTC CAC TAT TTC TTA TGT CAT TCG TTA 1865
 GTC TAG ATG TTT CTA AAC ATA TAA ATG TGA ATT TAA TCA ATT CCT TTC ATA GTT 1917
 TTA TAA TTC TCT GCG AGT TCC TTA TGA TAG AGT TTA TAA AAC AGT CCT GTG TAA 1971
 ACT GCT GGA AGT TCT TCC GCA ATT C

FIGURE 2
Human Acidic FGF

TGC ATT TTG TGC CTT TGC TGG AAG AAC CGA CTA CAG GTT TGT TCA ATT TCT TAC	27	54
AGT CTT GAA AGC GCC ACA AGC AGC AGC TGC TGA GCC MET Ala Glu Gly Glu Ile	81	108
ACC ACC TTC ACA GCC CTG ACC GAG AAG TTT AAT CTG CCT CCA GGG AAT TAC AAG	135	162
Thr Thr Phe Thr Ala Leu Thr Glu Lys Phe Asn Leu Pro Thr Gly Asn Tyr Lys	10	20
AAG CCC AAA CTC CTC TAC TGT AGC AAC GGG GGC CAC TTC CTG AGG ATC CTT CCG	189	216
Lys Pro Lys Leu Leu Tyr Cys Ser Asn Gly Gly His Phe Leu Arg Ile Leu Pro	30	40
GAT GGC ACA GTG GAT GGG ACA AGG GAC AGG AGC GAC CAG CAC ATT CAG CTG CAG	243	270
Asp Gly Thr Val Asp Gly Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln	50	60
CTC AGT GCG GAA AGC GTG GGG GAG GTG TAT ATA AAG AGT ACC GAG ACT GGC CAG	297	324
Leu Ser Ala Glu Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln	70	
TAC TTG GCC ATG GAC ACC GAC GGG CTT TTA TAC GGC TCA CAG ACA CCA AAT GAG	351	378
Tyr Leu Ala MET Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu	80	90
GAA TGT TTG TTC CTG GAA AGG CTG GAG GAG AAC CAT TAC AAC ACC TAT ATA TCC	405	432
Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr Ile Ser	100	110
AAG AAG CAT GCA GAG AAG AAT TGG TTT GTT GGC CTC AAG AAG AAT GGG AGC TGC	459	486
Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys Asn Gly Ser Cys	120	130
AAA CGC GGT CCT CGG ACT CAC TAT GGC CAG AAA GCA ATC TTG TTT CTC CCC CTG	513	540
Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala Ile Leu Phe Leu Pro Leu	140	150
CCA GTC TCT TCT GAT TAA AGA GAT CTG TTC TGG GTG TTG ACC ACT CCA GAG AAG	567	594
Pro Val Ser Ser Asp	155	
TTT CGA GGG GTC CTC ACC TGG TTG ACC CAA AAA TGT TCC CTT GA	621	

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FIGURE 5

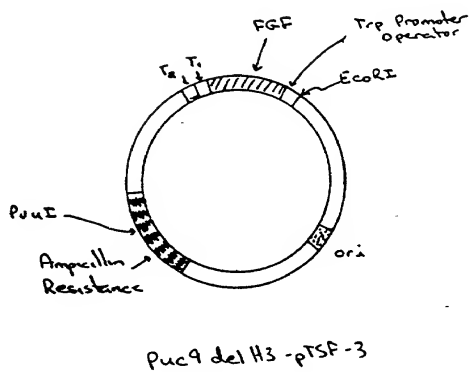
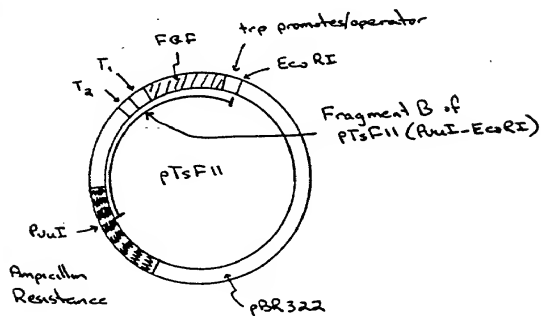
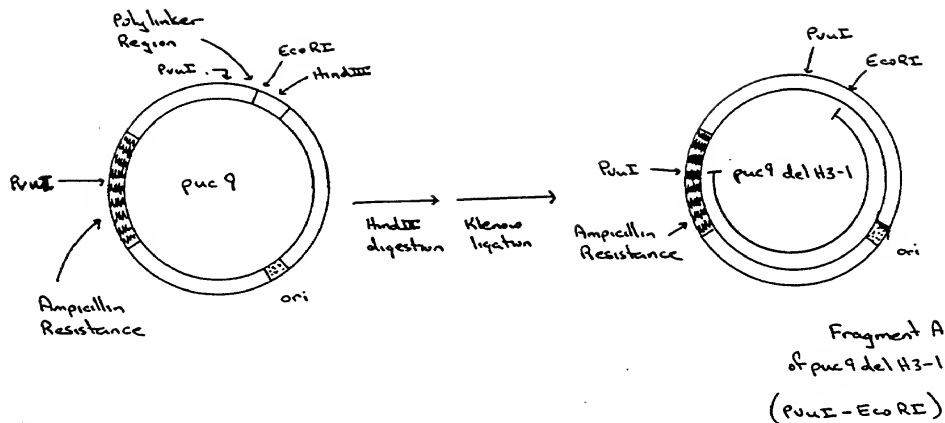


FIGURE 6

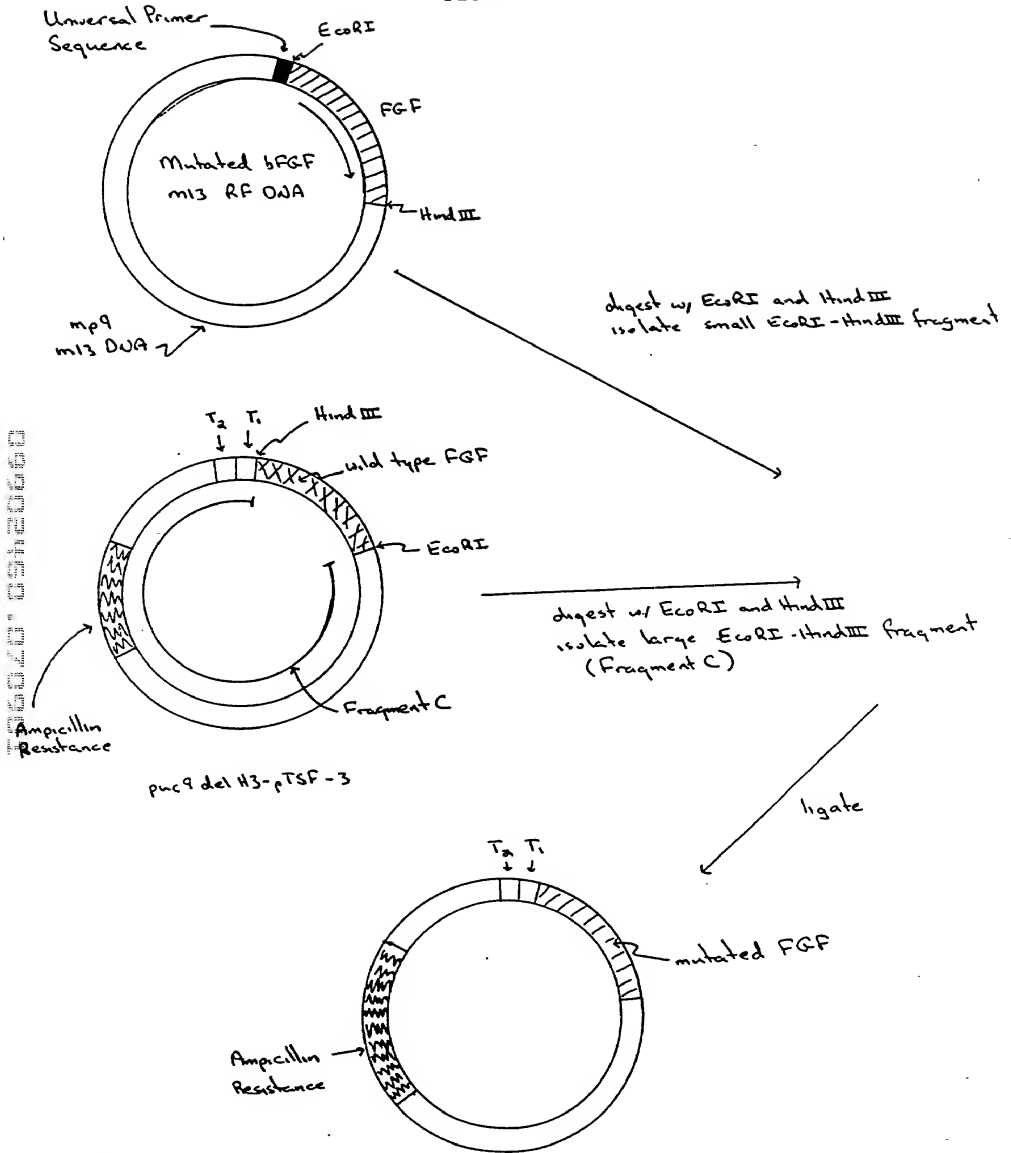


FIGURE 7

